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Thr Leu Ser Leu Thr Cys Ala Ile Ser Gly Gly Ser Ile Gly Ser Gly 20 25 30

Gly Ser Ile Arg Ser Thr Arg Trp Trp Ser Trp Val Arg Gln Ser Pro 35 40 45

Gly Lys Gly Leu Glu Trp Ile Gly Glu Ile Tyr His Ser Gly Ser Thr
50 55 60

Asn Tyr Asn Pro Ser Leu Lys Ser Arg Val Thr Ile Ser Leu Asp Lys 65 70 . 75 80

Ser Arg Asn His Phe Ser Leu Arg Leu Asn Ser Val Thr Ala Ala Asp 85 90 95

Thr Ala Val Tyr Tyr Cys Ala Arg Gln Asp Tyr Gly Asp Ser Gly Asp

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ceggggaagtt cececaceae tgtgatetat gaggataace aaagaceete tggggteeet 180

gateggttet etggeteeat egacagetee teeaactetg eeteceteae egtetetgga 240

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Thr Val Thr Ile Ser Cys Thr Arg Ser Ser Gly Ser Ile Ala Ser Asn 20 25 30

Ser Val Gln Trp Tyr Gln Gln Arg Pro Gly Ser Ser Pro Thr Thr Val 35 40 45

Ile Tyr Glu Asp Asn Gln Arg Pro Ser Gly Val Pro Asp Arg Phe Ser 50 55 60

Gly Ser Ile Asp Ser Ser Ser Asn Ser Ala Ser Leu Thr Val Ser Gly
65 70 75 80

Leu Lys Thr Glu Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Ser Asp Ser 85 90 95

Ser Ala Val Val Phe Gly Ser Gly Thr Lys Leu Thr Val Leu
100 105 110

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<212> DNA

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Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Arg Phe Thr Ser Tyr 20 25 30

Gly Ile Ser Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
35 40 45

Gly Trp Ile Ser Ala Tyr Asn Gly Asn Thr Asn Tyr Ala Gln Lys Leu 50 55 60

Gln Gly Arg Val Thr Met Thr Thr Asp Thr Ser Thr Asn Thr Ala Tyr
65 70 75 80

Met Glu Leu Arg Ser Leu Arg Ser Asp Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Arg Asp Ala Asp Tyr Ser Ser Gly Ser Gly Tyr Trp Gly Gln Gly
100 105 110

Thr Leu Val Thr Val Ser Ser 115

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<212> DNA

<213> Homo sapiens

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<213> Homo sapiens

<400> 8

Ser Tyr Glu Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly Gln

1 10 15

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20 25 30

Tyr Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Val Met Val Ile Tyr 35 40 45

Lys Asp Thr Glu Arg Pro Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser 50 55 60

Ser Ser Gly Thr Lys Val Thr Leu Thr Ile Ser Gly Val Gln Ala Glu
65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Ala Asp Asn Ser Ile Thr Tyr 85 90 95

Arg Val Phe Gly Gly Gly Thr Lys Val Thr Val Leu
100 105

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cctggacaag ggcttgagtg gatgggaata atcaacccta gaggtgccac cataagctac 180

gcacagaagt tccagggcag agtcaccatg accagggaca cgtccacgag tacagtctac 240

atggaactga gaaacttgaa atctgaggac acggccctgt attactgtgc tactgcaggc 300

atctatggtt ttgactttga ctactggggc agaggaaccc tggtcaccgt ctcctca 357

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<400> 10

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1 5 10 15

Ser Val Arg Val Ser Cys Lys Ala Ser Gly Tyr Thr Leu Thr Ser Tyr 20 25 30

Tyr Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met 35 40 45

Gly Ile Ile Asn Pro Arg Gly Ala Thr Ile Ser Tyr Ala Gln Lys Phe 50 55 60

Gln Gly Arg Val Thr Met Thr Arg Asp Thr Ser Thr Ser Thr Val Tyr
65 70 75 80

Met Glu Leu Arg Asn Leu Lys Ser Glu Asp Thr Ala Leu Tyr Tyr Cys 85 90 95

Ala Thr Ala Gly Ile Tyr Gly Phe Asp Phe Asp Tyr Trp Gly Arg Gly
100 105 110

Thr Leu Val Thr Val Ser Ser 115

<210> 11

<211> 333

<212> DNA

<213> Homo sapiens

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cacceaggea aageeeeaa acteateatt tatgatgtea etaaceggee etcaggggtt 180

tetgateget tetetggete caagtetgge aacaeggeet ecetgaeeat etetgggete 240

etggetgagg aegaggtga ttattaetge ageteataea caattgttae caatttegag 300

gttcttttcg gcggagggac caagctgacc gtc

333

<210> 12

<211> 111

<212> PRT

<213> Homo sapiens

<400> 12

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1 5 10 15

Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Asn Asp Val Gly Gly Tyr
20 25 30

Asn Tyr Val Ser Trp Tyr Gln His His Pro Gly Lys Ala Pro Lys Leu 35 40 45

Ile Ile Tyr Asp Val Thr Asn Arg Pro Ser Gly Val Ser Asp Arg Phe
50 55 60

Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Thr Ile Ser Gly Leu 65 70 75 80

Leu Ala Glu Asp Glu Gly Asp Tyr Tyr Cys Ser Ser Tyr Thr Ile Val 85 90 95

Thr Asn Phe Glu Val Leu Phe Gly Gly Gly Thr Lys Leu Thr Val

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Ala Tyr Tyr Trp Ser Trp Ile Arg Gln His Pro Gly Lys Gly Leu Glu

40

35

Trp Ile Gly Tyr Ile Tyr Tyr Asn Gly Asn Thr Tyr Tyr Asn Pro Ser 50 55 60

Leu Arg Ser Leu Val Thr Ile Ser Val Asp Ala Ser Lys Asn Gln Phe 65 70 75 80

Ser Leu Lys Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Val Tyr Tyr 85 90 95

Cys Ala Arg Ala Ser Asp Tyr Val Trp Gly Gly Tyr Arg Tyr Met Asp 100 105 110

Ala Phe Asp Ile Trp Gly Arg Gly Thr Leu Ile Thr Val Ser Ser 115 120 125

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<212> DNA

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ceaggaacgg cececaaact ceteatetat ggtaataate ageggeeete aggggteeet 180

gaeegattet etggeteeaa gtetggeaec teageeteee tggeeateag tgggeteeag 240

tetgagaatg aggetgatta ttaetgtgea geatgggatg acageetgaa tggteeggta 300

tteggeegag ggaeeaaggt caeegteeta ggtgag

<210> 16

<211> 112

<212> PRT

<213> Homo sapiens

<400> 16

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Arg Val Thr Ile Ser Cys Ser Gly Ser Asn Ser Asn Ile Gly Ser Asn 20 25 30

Ser Val Asn Trp Tyr Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu 35 40 45

Ile Tyr Gly Asn Asn Gln Arg Pro Ser Gly Val Pro Asp Arg Phe Ser 50 55 60

Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Leu Gln
65 70 75 80

Ser Glu Asn Glu Ala Asp Tyr Tyr Cys Ala Ala Trp Asp Asp Ser Leu 85 90 95

Asn Gly Pro Val Phe Gly Arg Gly Thr Lys Val Thr Val Leu Gly Glu
100 105 110

<210> 17

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Gln Asp Tyr Gly Asp Ser Gly Asp Trp Tyr Phe Asp Leu
                                   10
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Thr Arg Ser Ser Gly Ser Ile Ala Ser Asn Ser Val Gln
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Gln Ser Ser Asp Ser Ser Ala Val Val
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Ser Tyr Gly Ile Ser
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<212> PRT

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<213> Homo sapiens

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 Gln Ser Ala Asp Asn Ser Ile Thr Tyr Arg Val
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<211> 5
<212> PRT
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Ser Tyr Tyr Ile His
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<211> 17
<212> PRT
<213> Homo sapiens
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Gly
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Ala Gly Ile Tyr Gly Phe Asp Phe Asp Tyr
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<213> Homo sapiens
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Thr Gly Thr Ser Asn Asp Val Gly Gly Tyr Asn Tyr Val Ser
               5
                                  10
<210> 33
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<213> Homo sapiens
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Asp Val Thr Asn Arg Pro Ser
1
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<211> 12 <212> PRT

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 <213> Homo sapiens
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 Ser Gly Ala Tyr Tyr Trp Ser
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<211> 16
<212> PRT
<213> Homo sapiens
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Tyr Ile Tyr Tyr Asn Gly Asn Thr Tyr Tyr Asn Pro Ser Leu Arg Ser
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                                                       15
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<211> 17
<212> PRT
<213> Homo sapiens
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Ile

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 <213> Homo sapiens
 <400> 38
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 <213> Homo sapiens
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Gly Asn Asn Gln Arg Pro Ser
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<212> PRT
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<212> PRT

<213> Homo sapiens

<400> 41

Met Gln Ile Pro Gln Ala Pro Trp Pro Val Val Trp Ala Val Leu Gln

1 10 15

Leu Gly Trp Arg Pro Gly Trp Phe Leu Asp Ser Pro Asp Arg Pro Trp
20 25 30

Asn Pro Pro Thr Phe Phe Pro Ala Leu Leu Val Val Thr Glu Gly Asp

Asn Ala Thr Phe Thr Cys Ser Phe Ser Asn Thr Ser Glu Ser Phe Val 50 55 60

Leu Asn Trp Tyr Arg Met Ser Pro Ser Asn Gln Thr Asp Lys Leu Ala .
65 70 75 80

Ala Phe Pro Glu Asp Arg Ser Gln Pro Gly Gln Asp Cys Arg Phe Arg 85 90 95

Val Thr Gln Leu Pro Asn Gly Arg Asp Phe His Met Ser Val Val Arg

Ala Arg Arg Asn Asp Ser Gly Thr Tyr Leu Cys Gly Ala Ile Ser Leu 115 120 125

Ala Pro Lys Ala Gln Ile Lys Glu Ser Leu Arg Ala Glu Leu Arg Val

Thr Glu Arg Arg Ala Glu Val Pro Thr Ala His Pro Ser Pro Ser Pro 145 150 155 160

Arg Pro Ala Gly Gln Phe Gln Thr Leu Val Val Gly Val Val Gly Gly 165 170 175

Leu Leu Gly Ser Leu Val Leu Leu Val Trp Val Leu Ala Val Ile Cys
180 185 190

Ser Arg Ala Ala Arg Gly Thr Ile Gly Ala Arg Arg Thr Gly Gln Pro

Leu Lys Glu Asp Pro Ser Ala Val Pro Val Phe Ser Val Asp Tyr Gly
210 215 220

Glu Leu Asp Phe Gln Trp Arg Glu Lys Thr Pro Glu Pro Pro Val Pro 225 230 235 240

Cys Val Pro Glu Gln Thr Glu Tyr Ala Thr Ile Val Phe Pro Ser Gly
245 250 255

Met Gly Thr Ser Ser Pro Ala Arg Gly Ser Ala Asp Gly Pro Arg 260 265 270

Ser Ala Gln Pro Leu Arg Pro Glu Asp Gly His Cys Ser Trp Pro Leu 275 280 285

<210> 42

<211> 320

<212> DNA

<213> Homo sapiens

<400> 42

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cctggaaggc agatagcagc cccgtcaagg cgggagtgga gaccaccaca ccctccaaac 180
aaagcaacaa caagtacgcg gccagcagct atctgagcct gacgcctgag cagtggaagt 240
cccacagaag ctacagctgc caggtcacgc atgaagggag caccgtggag aagacagtgg 300
cccctacaga atgttcatag

<210> 43

<211> 106

<212> PRT

<213> Homo sapiens

<400> 43

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Glu Glu Leu Gln Ala Asn Lys Ala Thr Leu Val Cys Leu Ile Ser Asp 20 25 30

Phe Tyr Pro Gly Ala Val Thr Val Ala Trp Lys Ala Asp Ser Ser Pro 35 40 45

Val Lys Ala Gly Val Glu Thr Thr Pro Ser Lys Gln Ser Asn Asn 50 55 60

Lys Tyr Ala Ala Ser Ser Tyr Leu Ser Leu Thr Pro Glu Gln Trp Lys 65 70 75 80

Ser His Arg Ser Tyr Ser Cys Gln Val Thr His Glu Gly Ser Thr Val 85 90 95

Glu Lys Thr Val Ala Pro Thr Glu Cys Ser 100 105

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<211> 960

<212> DNA

<213> Homo sapiens

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ggaacteagg egecetgace ageggegtge acacettece ggetgteeta cagteeteag 180
gactetacte ecteageage gtggtgaceg tgeeeteag eagettggge acceagacet 240
acatetgeaa egtgaateae aageeeagea acaceaaggt ggacaagaaa gttgageeca 300
aatettgtga caaaacteae acatgeeeae egtgeeeage acetgaacte etggggggae 360
egteagtett ectetteeee ecaaaaccea aggacaceet eatgatetee eggaceeetg 420
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gcacgtaceg tgtggteage gteeteaceg teetgcacea ggactggetg aatggcaagg 600

agtacaagtg caaggtetec aacaaagcec teccageece categagaaa accateteca 660 aageeaaagg geageeega gaaceacagg tgtacaceet geeeceatee egggaggaga 720 tgaccaagaa ceaggteage etgacetgee tggteaaagg ettetateee agegacateeg 780 eegtggagtg ggagageaat gggeageegg agaacaacta caagaceaeg eetecegtge 840 tggacteega eggeteette tteetetata geaageteae egtggacaag ageaggtgge 900 ageaggggaa egtettetea tgeteegtga tgeatgagge tetgcacaac cactacaege 960

<210> 45

<211> 330

<212> PRT

<213> Homo sapiens

<400> 45

Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys

1 10 15

Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr 20 25 30

Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser 35 40 45

Gly Val His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser 50 55 60

Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr 65 70 75 80

Tyr Ile Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys 85 90 95

Lys Val Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys
100 105 110

Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro 115 120 125

Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys
130 135 140

Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp 145 150 155 160

Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu 165 170 175

Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu 180 185 190

His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn 195 200 205

Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly
210 220

Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu 225 230 235 240

Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr 245 250 255

Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn 260 265 270

Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe 275 280 285

Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn 290 295 300

Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr 305 310 315 320

Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys 325 330

<210> 46

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<212> DNA

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<400> 46

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gtccgccagg ctccagggaa ggggctggag tgggtctcag ctattagtgg tagtggtggt 180

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<210> 47

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Phe Ser Ser Tyr Trp Cys Asp Arg Met Ser Trp Val Arg Gln Ala Pro
35 40 45

Gly Lys Gly Leu Glu Trp Val Ser Ala Ile Ser Gly Ser Gly Gly Ser 50 55 60

Thr Tyr Tyr Ala Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp 65 70 75 80

Asn Ser Lys Asn Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu 85 90 95

Asp Thr Ala Val Tyr Tyr Cys Ala Lys Glu Asn Trp Gly Ser Tyr Phe
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Asp Leu Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser

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cagcagaaac cagggagage cectaaggte ttgatetata aggeatetae tttagaaagt 180
ggggteecat caaggtteag eggeagtgga tetgggacag attteactet caccateage 240
agtetgeaac etgaagattt tgeaacttae taetgteaac agagttacag taeceegtgg 300
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<210> 49

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<213> Homo sapiens

<400> 49

Gly Val His Ser Asp Ile Val Met Thr Gln Ser Pro Ser Thr Leu Ser 1 5 10 15

Ala Ser Val Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Gly
20 25 30

Ile Ser Ser Trp Leu Ala Trp Tyr Gln Gln Lys Pro Gly Arg Ala Pro 35 40 45

Lys Val Leu Ile Tyr Lys Ala Ser Thr Leu Glu Ser Gly Val Pro Ser 50 55 60

Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser 65 70 75 80

Ser Leu Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr 85 90 95

Ser Thr Pro Trp Thr Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys Arg

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<210> 51

<211> 17

<212> PRT

<213> Homo sapiens

<400> 51

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Ala Ile Ser Gly Ser Gly Ser Thr Tyr Tyr Ala Asp Ser Val Lys
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 <210> 52
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 <400> 52
Glu Asn Trp Gly Ser Tyr Phe Asp Leu
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Asn Ala Thr Phe Thr Cys Ser Leu Ser Asn Trp Ser Glu Asp Leu Met 50 55 60

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Ala Phe Cys Asn Gly Leu Ser Gln Pro Val Gln Asp Ala Arg Phe Gln 85 90 95

Ile Ile Gln Leu Pro Asn Arg His Asp Phe His Met Asn Ile Leu Asp

Thr Arg Arg Asn Asp Ser Gly Ile Tyr Leu Cys Gly Ala Ile Ser Leu 115 120 125

His Pro Lys Ala Lys Ile Glu Glu Ser Pro Gly Ala Glu Leu Val Val
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Thr Glu Arg Ile Leu Glu Thr Ser Thr Arg Tyr Pro Ser Pro Ser Pro 145

Lys Pro Glu Gly Arg Phe Gln Gly Met Val Ile Gly Ile Met Ser Ala
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Leu Val Gly Ile Pro Val Leu Leu Leu Leu Ala Trp Ala Leu Ala Val

Phe Cys Ser Thr Ser Met Ser Glu Ala Arg Gly Ala Gly Ser Lys Asp

Asp Thr Leu Lys Glu Glu Pro Ser Ala Ala Pro Val Pro Ser Val Ala 210 215 220

Tyr Glu Glu Leu Asp Phe Gln Gly Arg Glu Lys Thr Pro Glu Leu Pro 225 230 235 240

Thr Ala Cys Val His Thr Glu Tyr Ala Thr Ile Val Phe Thr Glu Gly
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Gln Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp 50 55 60

Ser Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr 65 70 75 80

Glu Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser 85 90 95

Ser Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys 100 105